RAW SEQUENCE LISTING PATENT APPLICATION US/08/572,027B

DATE: 05/06/98 TIME: 12:59:02

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This Raw Listing contains the General Information Section and up to the first Wages

SEQUENCE LISTING 1 2 General Information: 3 (1)5 (i) APPLICANT: DeBonte, L. et al. 6 (ii) TITLE OF INVENTION: PLANTS HAVING MUTANT SEQUENCES THAT CONFER 7 8 ALTERED FATTY ACID PROFILES 9 10 (iii) NUMBER OF SEQUENCES: 8 11 12 (iv) CORRESPONDENCE ADDRESS: 13 14 (A) ADDRESSEE: Fish & Richardson P.C., P.A. 15 (B) STREET: 60 South Sixth Street, Suite 3300 16 (C) CITY: Minneapolis (D) STATE: MN 17 18 (E) COUNTRY: USA (F) ZIP: 55402 19 20 21 (v) COMPUTER READABLE FORM: 22 (A) MEDIUM TYPE: Diskette 23 (B) COMPUTER: IBM compatible 24 (C) OPERATING SYSTEM: Windows95 25 (D) SOFTWARE: FastSEQ for Windows Version 2.0 26 27 (vi) CURRENT APPLICATION DATA: 28 (A) APPLICATION NUMBER: US 08/572,027 29 (B) FILING DATE: 14-DEC-1995 30 (C) CLASSIFICATION: 31 32 (Vi) PRIOR APPLICATION DATA: 33 (A) APPLICATION NUMBER: US 08/416,497 (B) FILING DATE: 04-APR-1995 34 35 (C) CLASSIFICATION: 36 37 (vi) PRIOR APPLICATION DATA: 38 (A) APPLICATION NUMBER: US 08/170,886 39 (B) FILING DATE: 21-DEC-1993 40 (C) CLASSIFICATION: 41 42 (vi) PRIOR APPLICATION DATA: 43 (A) APPLICATION NUMBER: US 07/739,965 RECEIA DCATE: 05-AUG-1991 44 45 46

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RAW SEQUENCE LISTING PATENT APPLICATION US/08/572,027B

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RAW SEQUENCE LISTING PATENT APPLICATION US/08/572,027B

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105									Tyr								
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107																	
108	CTA	ACC	GGC	GTC	TGG	GTC	ATA	GCC	CAC	GAA	TGC	GGC	CAC	CAC	GCC	TTC	336
109	Leu	Thr	Gly	Val	Trp	Val	Ile	Ala	His	Glu	Cys	Gly	His	His	Ala	Phe	
110				100					105					110			
111																	
112									ACC								384
113	Ser	Asp	Tyr	Gln	Trp	Leu	Asp	_	Thr	Val	Gly	Leu		Phe	His	Ser	
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121									Arg								400
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123		AAG	TCA	GAC	ATC	_	TGG	TAC	GGC	AAG		CTC	AAC	AAC	CCT		528
124									Gly								
125	1	-		-	165	-	-	-	-	170	-				175		
126																	
127	GGA	CGC	ACC	GTG	ATG	TTA	ACG	GTT	CAG	TTC	ACT	CTC	GGC	TGG	CCG	TTG	576
128	Gly	Arg	Thr	Val	Met	Leu	Thr	Val	Gln	Phe	Thr	Leu	Gly	Trp	Pro	Leu	
129				180					185					190			
130																	
131									AGA								624
132	Tyr	Leu		Pne	Asn	vaı	ser	_	Arg	Pro	туг	Asp		GTÀ	Pne	Arg	
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139	CAG	ATA	TAC	ATC	TCC	GAC	GCT	GGC	ATC	CTC	GCC	GTC	TGC	TAC	GGT	CTC	720
140	Gln	Ile	Tyr	Ile	Ser	Asp	Ala	Gly	Ile	Leu	Ala	Val	Cys	Tyr	Gly	Leu	
141	225		_			230		_			235					240	
142																	
143									GTG								768
144	Phe	Arg	Tyr	Ala		Gly	Gln	Gly	Val		Ser	Met	Val	Cys		Tyr	
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RAW SEQUENCE LISTING PATENT APPLICATION US/08/572,027B

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ASP Trp Phe Arg Gly Ala Leu Ala Thr Val Asp Arg Asp Tyr Gly Ile 157 290 295 300	154																		
158	155	GAT	TGG	TTC	AGG	GGA	GCT	TTG	GCT	ACC	GTT	GAC	AGA	GAC	TAC	GGA	ATC		912
158	156	Asp	Trp	Phe	Arg	Gly	Ala	Leu	Ala	Thr	Val	Asp	Arg	Asp	Tyr	Gly	Ile		
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170 171 CTT AAG CCG ATG TGG AGG GAG GCG AAG GAG TGT ATC TAT GTG GAA CCG 172 Val Lys Ala Met Trp Arg Glu Ala Lys Glu Cys Ile Tyr Val Glu Pro 173 355 360 365 174 175 GAC AGG CAA GGT GAG AAG AAA GGT GTG TTC TGG TAC AAC AAT AAG TTA T 176 Asp Arg Gln Gly Glu Lys Lys Gly Val Phe Trp Tyr Asn Asn Lys Leu 177 370 375 380 1159 180 181 179 GA	168	Ile	Lys	Pro	Ile	Leu	Gly	Glu	Tyr	Tyr	Gln	Phe	Asp	Gly	Thr	Pro	Val		
171 GTT AAG GCG ATG TGG AGG GAG GCG AAG GAG TGT ATC TAT GTG GAA CCG 1104 172 173 174 175 185	169				340					345					350				
172 Val Lys Ala Met Trp Arg Glu Ala Lys Glu Cys Ile Tyr Val Glu Pro 355 360 365	170																		
173	171	GTT	AAG	GCG	ATG	TGG	AGG	GAG	GCG	AAG	GAG	TGT	ATC	TAT	GTG	GAA	CCG		1104
173	172	Val	Lys	Ala	Met	Trp	Arq	Glu	Ala	Lys	Glu	Cys	Ile	Tyr	Val	Glu	Pro		
174 175 GAC AGG CAA GGT GAG AAG AAA GGT GTG TTC TGG TAC AAC AAT AAG TTA T 176 Asp Arg Gln Gly Glu Lys Lys Gly Val Phe Trp Tyr Asn Asn Lys Leu 177 178 179 GA 179 GA 180 181 182 (2) INFORMATION FOR SEQ ID NO:2: 183 184 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 384 amino acids 186 (B) TyPE: amino acid 187 (D) TOPOLOGY: linear 188 189 (ii) MOLECULE TYPE: protein 190 191 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2: 192 193 Met Gly Ala Gly Gly Arg Met Gln Val Ser Pro Pro Ser Lys Lys Ser 194 1 5 10 15 195 196 Glu Thr Asp Thr Ile Lys Arg Val Pro Cys Glu Thr Pro Pro Phe Thr 197 20 25 30 198 199 Val Gly Glu Leu Lys Lys Ala Ile Pro Pro His Cys Phe Lys Arg Ser 200 35 40 201 202 21e Pro Arg Ser Phe Ser Tyr Leu Ile Trp Asp Ile Ile Ile Ala Ser 203 204 204	173		-			-				-		-		_					
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194	192																		
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204		тте		arg	ser	rne	ser	_	ьeu	тте	тгр	ASP		тте	тте	Ald	ser		
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205 Cys Pne Tyr Tyr Xaa Ala Thr Thr Tyr Phe Pro Leu Leu Pro His Pro			_,	_	_			,	_,		1	_	_	_	_		_		
	205	Cys	Phe	Tyr	Tyr	хаа	Ala	Thr	Thr	Tyr	Phe	Pro	Leu	Leu	Pro	His	Pro		

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210		_	_	_		_						_				
211 212 213	Leu	Thr	Gly	Val 100	Trp	Val	Ile	Ala	His 105	Glu	Cys	Gly	His	His 110	Ala	Phe
214 215	Ser	Asp	Tyr 115	Gln	Trp	Leu	Asp	Asp 120	Thr	Val	Gly	Leu	Ile 125	Phe	His	Ser
216 217 218	Phe	Leu 130	Leu	Val	Pro	Tyr	Phe 135	Ser	Trp	Lys	Tyr	Ser 140	His	Arg	Ser	His
219 220 221 222	His 145	Ser	Asn	Thr	Gly	Ser 150	Leu	Glu	Arg	Asp	Glu 155	Val	Phe	Val	Pro	Lys 160
223 224	Lys	Lys	Ser	Asp	Ile 165	Lys	Trp	Tyr	Gly	Lys 170	туr	Leu	Asn	Asn	Pro 175	Leu
225 226 227 228	Gly	Arg	Thr	Val 180	Met	Leu	Thr	Val	Gln 185	Phe	Thr	Leu	Gly	Trp 190	Pro	Leu
229 230 231	Tyr	Leu	Ala 195	Phe	Asn	Val	Ser	Gly 200	Arg	Pro	Туr	Asp	Gly 205	Gly	Phe	Arg
232 233 234	Cys	His 210	Phe	His	Pro	Asn	Ala 215	Pro	Ile	Tyr	Asn	Asp 220	Arg	Glu	Arg	Leu
235 236 237	Gln 225	Ile	Tyr	Ile	Ser	Asp 230	Ala	Gly	Ile	Leu	Ala 235	Val	Cys	Туr	Gly	Leu 240
238 239 240	Phe	Arg	Tyr	Ala	Ala 245	Gly	Gln	Gly	Val	Ala 250	Ser	Met	Val	Cys	Phe 255	Tyr
241 242 243	Gly	Val	Pro	Leu 260	Leu	Ile	Val	Asn	Gly 265	Phe	Leu	Val	Leu	11e 270	Thr	Tyr
244 245 246	Leu	Gln	His 275	Thr	His	Pro	Ser	Leu 280	Pro	His	Tyr	Asp	Ser 285	Ser	Glu	Trp
247 248 249	Asp	Trp 290	Phe	Arg	Gly	Ala	Leu 295	Ala	Thr	Val	Asp	Arg 300	Asp	Tyr	Gly	Ile
250 251 252	Leu 305	Asn	Lys	Val	Phe	His 310	Asn	Ile	Thr	Asp	Thr 315	His	Val	Ala	His	His 320
252 253 254 255	Pro	Phe	Ser	Thr	Met 325	Pro	His	Tyr	His	Al a 330	Met	Glu	Ala	Thr	Lys 335	Ala
256 257 258	Ile	Lys	Pro	Ile 340	Leu	Gly	Glu	Tyr	Tyr 345	Gln	Phe	Asp	Gly	Thr 350	Pro	Val

SEQUENCE VERIFICATION REPORT PATENT APPLICATION US/08/572,027B

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